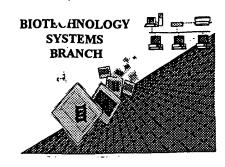
0905

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/930,9/5Source: 0/966Date Processed by STIC: 8/23/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

DATE: 08/23/2001

OIPE

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PATENT APPLICATION: US/09/930,915
                                                              TIME: 17:29:04
                                                                            Does Not Comply
                     Input Set : A:\Icc-1022.app
                                                                       Corrected Diskette Needed
                     Output Set: N:\CRF3\08162001\I930915.raw
      3 <110> APPLICANT: Birkett, Ashley J.
      5 <120> TITLE OF INVENTION: IMMUNOGENIC HBc CHIMER PARTICLES HAVING ENHANCED
      6
              STABILITY
      8 <130> FILE REFERENCE: 4564/81175 ICC-102.2
 H-> 10 <140> CURRENT APPLICATION NUMBER: US/09/930,915
     11 <141> CURRENT FILING DATE: 2001-08-15
     13 <150> PRIOR APPLICATION NUMBER: 60/226,867
     14 <151> PRIOR FILING DATE: 2000-08-22
     16 <150> PRIOR APPLICATION NUMBER: 60/225,843
     17 <151> PRIOR FILING DATE: 2000-08-16
     19 <160> NUMBER OF SEQ ID NOS: 313
     21 <170> SOFTWARE: PatentIn Ver. 2.1
ERRORED SEQUENCES
     2140 <210> SEQ ID NO: 188
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     2142 <212> TYPE: PRT
     2143 <213> ORGANISM: Hepatitis C virus
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     2218 <213> ORGANISM: Plasmodium vivax
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2222 1
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     4033 <212> TYPE: DNA
     4034 <213> ORGANISM: Hepatitis B virus
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     4039 geettagagt etectgagea ttgeteacet eaceatactg caeteaggea agecattete 180
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4040 tgctgggggg aattgatgac tctagctacc tgggtgggta ataatttgca agatccagca 240

RAW SEQUENCE LISTING

RAW SEQUENCE LISTING

DATE: 08/23/2001

PATENT APPLICATION: US/09/930,915

TIME: 17:29:05

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Output Set: N:\CRF3\08162001\I930915.raw

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	4044	tcaacacttc	cggaaactac	tgttgttaga	cgacgggacc	gaggcaggtc	ccctagaaga	480
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	4042	ttgtggtttc	atatatcttg	ccttacttt	ggaagagaga	ctgtacttga	atatttggtc	360
		, ,	, ,	_	, ,	gtttaaagat	J J	





VERIFICATION SUMMARY

PATENT APPLICATION: US/09/930,915

DATE: 08/23/2001 TIME: 17:29:06

Input Set : A:\Icc-1022.app

Output Set: N:\CRF3\08162001\1930915.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number

L:2149 M:252 E: No. of Seq. differs, <211>LENGTH:Input:16 Found:17 SEQ:188 L:2211 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:193 L:2221 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:194

L:3325 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:258

L:3411 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:260

L:4046 M:252 E: No. of Seq. differs, <211>LENGTH:Input:554 Found:555 SEQ:276